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## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

<b>(51) International Patent Classification <sup>4</sup> :</b>  <b>A61K 37/42, C07K 7/18</b>	<b>A1</b>	<b>(11) International Publication Number:</b> <b>WO 89/ 01781</b>  <b>(43) International Publication Date:</b> <b>9 March 1989 (09.03.89)</b>
<b>(21) International Application Number:</b> <b>PCT/US88/02960</b>  <b>(22) International Filing Date:</b> <b>29 August 1988 (29.08.88)</b>  <b>(31) Priority Application Number:</b> <b>091,995</b>  <b>(32) Priority Date:</b> <b>2 September 1987 (02.09.87)</b>  <b>(33) Priority Country:</b> <b>US</b>  <b>(71)(72) Applicants and Inventors:</b> <b>STEWART, John, M. [US/US]; 3690 East Dartmouth Avenue, Denver, CO 80210 (US). VAVREK, Raymond, J. [US/US]; 1250 Poplar Street, Denver, CO 80220 (US).</b>  <b>(74) Agents:</b> <b>BRENEMAN, William, D. et al.; Breneman &amp; Georges, Suite 290 - International Square, 1850 K Street, N.W., Washington, DC 20006 (US).</b>		<b>(81) Designated States:</b> <b>AT (European patent), AU, BE (European patent), CH (European patent), DE (European patent), DK, FI, FR (European patent), GB (European patent), IT (European patent), JP, KR, LU (European patent), NL (European patent), NO, SE (European patent).</b>  <b>Published</b> <i>With international search report.          Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.</i>
<b>(54) Title:</b> <b>BRADYKININ ANTAGONIST PEPTIDES</b>  <b>(57) Abstract</b>  <p>The substitution of the L-Pro. at the 7-position of the peptide hormone bradykinin or other substituted analogs of bradykinin with an aliphatic, cyclic or aromatic amino acid of the D-configuration converts bradykinin agonists into a bradykinin antagonist. The invention further includes additional modifications at other positions within the novel 7-position modified bradykinin antagonists including C-terminal and N-terminal extensions and replacement of arginine in the one- and/or two positions which increase enzyme resistance, antagonist potency and/or specificity of the new bradykinin antagonists. The analogs produced are useful in treating conditions and diseases of the mammal and human body in which an excess of bradykinin or related kinins are produced or injected as by bites into the body.</p>		

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## BRADYKININ ANTAGONIST PEPTIDES

## BACKGROUND OF THE INVENTION

1. Cross Reference To Related Applications

The invention pertains to new and useful bradykinin antagonist peptides and is related to the subject matter of U.S. Application Serial No. \_\_\_\_\_ filed concurrently herewith.

2. Field Of The Invention

The invention relates to novel biologically active peptides which act as antagonists of the biological activities of bradykinin, their pharmaceutically acceptable salts, and their application as therapeutic agents. More particularly the invention pertains to the substitution of L-Pro at position 7 to convert bradykinin agonists into antagonists together with other modifications and substitutions at other positions including C and N-terminal extensions and modifications, amino acid deletions and the substitution of arginine in the one and nine positions of the novel bradykinin antagonists.

3. Description Of The Prior Art

In the 25 years since the sequence of the potent mammalian vasodilator peptide bradykinin was described and synthesized (Boissonnas et al., Experientia 16: 326, 1960).

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several hundred sequence-related peptide analogs have been synthesized and assayed in biological systems (Schroeder, in Handbook of Experimental Pharmacology, Vol. 25, (Springer Verlag) pp. 324-350, 1970) (Stewart, Handbook of Experimental Pharmacology, Vol. 25 (Supplement), (Springer Verlag) pp. 227-272, 1979). The objective in these studies was to investigate the varied physiological and pharmacological roles of bradykinin.

Bradykinin, and its physiologically important related peptides kallidin (Lys-bradykinin) and Met-Lys-bradykinin, exhibit physiological actions which qualify them as mediators of inflammatory reactions, hypotensive states, and pain. Bradykinin is overproduced in pathological conditions such as septic shock (Robinson et al., Am. J. Med. 59:61, 1975) and hemorrhagic (Hirsch et al., J. Surg. Res. 17:147, 1974), anaphylaxis, (Collier and James, J. Physiol. 160:15P, 1966), arthritis (Jasani et al., Am. Rheum. Dis. 28:497, 1969; Hamberg et al., Agents Actions 8:50, 1978; Sharma et al., Arch. Int. Pharmacodyn 262:279, 1983), rhinitis (Proud et al., J. Clin. Invest. 72:1678, 1983; Naclerio et al., Clin. Res. 33:613A, 1985), asthma (Christiansen et al., J. Clin. Invest. 79:188 - 197, 1987), inflammatory bowel disease (Zeitlin and Smith, Gut 14:133 - 138, 1973), and certain other conditions including acute pancreatitis, post-gastrectomy dumping syndrome,

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carcinoid syndrome, migraine, and angioneurotic edema (Leme, Handb. Exp. Pharmacol. 50/I:464 - 522, 1978). The production of bradykinin from the plasma results in pain at the site of the pathological condition, and the overproduction intensifies the pain directly or via stimulation by bradykinin of the activation of the arachidonic acid pathway which produces prostaglandins and leukotrienes, the more distal and actual mediators of inflammation. Literature references describing these actions of bradykinin and related peptides are found in Handbook of Experimental Pharmacology, Vol. 25, Springer-Verlag, 1970 and Vol. 25 Supplement, 1979.

Bradykinin as discussed has been found to be produced in inflammatory reactions in the intestine provoking contraction of smooth muscle and secretion of fluid and ions. The existence of specific bradykinin receptors in the mucosal lining of the intestine and intestinal smooth muscle is demonstrated by Manning, et al in Nature (229: 256-259, 1982) showing the influence of bradykinin in very low concentrations upon fluid and ion secretion.

The production of bradykinin and associated pain in angina has been studied and reported by Kimura, et al in American Heart Journal (85: 635-647, 1973) and by Staszewska - Barczak, et al in Cardiovascular Research (10: 314-327, 1976). The reported action of bradykinin and prostaglandins acting in

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concert are the natural stimulus for excitation of the sensory receptors signalling the pain of myocardial ischaemia.

Bradykinin and bradykinin - related kinins are not only produced by the animal but may also be injected as a result of stings and bites. It is known that insects such as hornets and wasps inject bradykinin related peptides which also cause pain, swelling and inflammation.

The search for understanding of the mechanism of action of bradykinin, which is essential for the development of useful tools for diagnostic use, and for the development of therapeutic agents aimed at alleviating the intense pain caused by the production and overproduction of bradykinin, has been severely hindered by the lack of specific sequence-related competitive antagonists of bradykinin.

Several non-peptide, non-specific and non-selective antagonists of one or more of the biological activities of bradykinin have been described among compounds as diverse as analgesics and anti-inflammatory substances, which act via the prostaglandin system and not directly on bradykinin biological receptors (Rocha e Silva and Leme, Med. Exp, 8:287, 1963). These are antihistamines (Gecse et al, J. Pharm. Pharmacol. 21: 544, 1969); bradykinin-antibodies (Grez et al, Eu. J. Pharmacol. 29: 35, 1974); benzodiazepine derivatives (Leme and Rocha Silva, Br. J. Pharmacol. 25: 50, 1965); high molecular

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weight ethylene oxide polymers (Wilkins and Back, Arch. Intl. Pharmacodynam. 209: 305, 1974); gallic acid esters (Posati et al., J. Agri. Food Chem. 18: 632, 1970) and serotonin inhibitors (Gomazkon and Shimkovich, Bull. Exptl. Biol. Med. 80: 6, 1975). None of these individual compounds or classes of compounds specifically inhibit bradykinin.

Heptyl esters of various amino acid-containing substances, such as single basic amino acids (ie. Arg and Lys) (Gecse, Adv. Exptl. Biol. Med. 70: 5, 1976), the dipeptide Phe-Gly (Gecse et al, Int. Aech. Allergy 41: 174, 1971), and of analogs of C- terminal peptide fragments of bradykinin (ie, Pro-Phe-Arg) (Claesson et al., Adv. Exptl. Med. Biol. 120B: 691, 1979) have been reported as anti-bradykinin substances. When tested in bradykinin assay systems they prove to be weak partial agonists/antagonists, depending on the dose, with little specificity for inhibiting bradykinin action.

Preparations of damaged vascular tissue have been reported to respond to bradykinin analogs which lack the C-terminal Arg residue, but not to bradykinin itself, and analogs of these des-Arg<sup>9</sup>-bradykinins have been developed as antagonists of this non-physiological activity of bradykinin. These antagonists have no significant bradykinin-like agonist effects, nor any antagonist effect on any of the physiologically significant kinin-responding systems (Regoli



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and Barabe, Pharmacol. Revs. 32:1,1980).

Several bradykinin analogs containing the O-methyl ether of Tyr residues at positions 5 and/or 8 have been reported to produce mixed agonist/antagonist activity on isolated uteri of galactosemic rats, but not on normal rats. The antagonism was not reliably reproducible in these animals (Stewart and Woolley, in Hypotensive Peptides, Springer Verlag, pp. 23-33, 1966).

Other changes in the bradykinin molecule have been additions of amino acids at the N-terminal end which affect the rate of enzymatic degradation of bradykinin in vivo.

The half life of bradykinin in the systemic circulation is less than 30 seconds (S.H. Ferreira & J.R. Vane, Br. J. Pharmacol. Chemotherap. 30:417, 1967). Bradykinin is completely destroyed (98-99% destruction) on a single passage through the pulmonary circulation (J. Roblero, J.W. Ryan and J.M. Stewart, Res. Commun. Pathol. Pharmacol. 6:207, 1973) as determined in the anesthetized rat by measuring the depressor effects of an agonist following intra-aortic (IA) (bypassing the pulmonary circulation) and intravenous (IV) administration. Resistance of bradykinin agonists to pulmonary kininase destruction in vivo is promoted by addition of single (ie, DArg-, DLys-, Lys-) and double (DLys-Lys-) basic amino acid residues to the N-terminal of the bradykinin sequence. The

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addition of the dipeptide Lys-Lys to the N-terminal of bradykinin agonists confers complete resistance to in vivo destruction on initial passage through the pulmonary circulation (Roblero, Ryan and Stewart, Res. Comm. Pathol. Pharmacol. 6:207, 1973).

#### SUMMARY OF INVENTION

The invention relates to the modification of the sequence of the mammalian peptide hormone bradykinin (Arg-Pro-Pro-Gly-Phe-Ser-Pro-Phe-Arg) and pharmaceutically acceptable salts thereof, at the Pro residue at position 7 in a unique manner which, for the first time, produces sequence-related analogues that act as specific and competitive inhibitors of the biological activities of bradykinin. The invention specifically relates to the substitution of the L-Pro at position 7 with substituted and unsubstituted aromatic amino acids of the D-configuration, a change which converts bradykinin agonists into antagonists, and includes additional modifications at other positions within the 7-position modified bradykinin antagonist which confer increased antagonist potency, resistance to enzymatic degradation and/or tissue specificity on the D-amino acid-containing bradykinin sequence. The invention further includes the necessary substitution of L-Pro at position 7 with substituted and unsubstituted amino acids of the D-configuration together with the substitution of

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arginine in the one and nine positions with D or L-cyclic (heterocyclic or alicyclic) amino acid residue, D or L aliphatic amino acid residue or a D or L substituted or unsubstituted aromatic acid residue. The invention also includes C-terminal modifications and extensions. More specifically, the invention relates to the peptides of the general formula:

Formula I

N-A1-B-C-D-W-X-Y-Z-A9-Cn  
(0 1 2 3 4 5 6 7 8 9 10 ) (position number)

Wherein N is a hydrogen atom or single acidic, basic or neutral aromatic amino acid residue of the D- or L- configuration, such as D-Arg, D-Lys or L-Thi, an N-terminal enzyme protecting group selected from the group comprising acyl-type protecting groups, aromatic urethane-type protecting groups, alkyl-type protecting groups, or alternately N is a di- or poly-peptide containing amino acids of the D- or L- configuration, such as Lys-Lys, Met-Lys, or Gly-Arg-Met-Lys;

A1 and A9 are either or both an Arg residue or other cyclic (heterocyclic or alicyclic) amino acid residue, aliphatic amino acid residue or an aromatic or substituted aromatic amino acid residue of the D or L configuration;

B is D- or L-pro residue, or other D- or L-cyclic (heterocyclic or alicyclic) or noncyclic aliphatic amino acid

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residue, such as L-hydroxyproline or a D or L-aromatic or substituted aromatic amino acid residue;

C is D- or L-Pro residue, or other cyclic (heterocyclic or alicyclic), aliphatic, aromatic or substituted aromatic amino acid residue of the D- or L-configuration;

D is a Gly residue or other aliphatic, cyclic, aromatic or substituted aromatic amino acid residue of the D or L-configuration, such as Ala;

W is a Phe residue of the D or L-configuration, or a substituted Phe or other aliphatic or aromatic amino acid residue of the D- or L- configuration, such as Leu, beta-2-thienyl-alanine (Thi) or 2-pyridyl-alanine (Pal) or a cyclic amino acid such as a D or L-Pro residue;

X is a Ser residue of the D- or L-configuration, a Gly residue, or other D- or L-aliphatic, cyclic or aromatic or substituted aromatic amino acid residue, such as pCl-D-Phe or D-Phe;

Y is a D- aromatic amino acid residue, or substituted aromatic amino acid residue, such as D-Phe, beta-(2-thienyl)-D Ala (DThi), beta-(2-pyridyl)-D-Ala (D-Pal),  $\beta$ -2-naphthyl-D Ala (D-Nal), DHis, D-homo-Phe (DhPhe), O-methyl-DTyr (DOMT), D-alphaphenyl-Gly (DPHg), DTrp, DTyr or pCl-DPhe (CDF);

Z is a Phe residue of the D or L configuration, or a

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substituted Phe or other aliphatic or aromatic amino acid residue of the D- or L-configuration, such as Leu, Thi or Pal or a cyclic amino acid such as a D or L-Pro; and

Cn is a hydroxyl group or a C-terminal extension such as an amide, alkoxy group, an acidic, basic or neutral aliphatic aromatic, or cyclic amino acid residue of the D- or L-configuration or a peptide extension composed of D- or L-amino acids.

In a preferred compound of the general formula I the substituents have the following identity: A = H, B = Pro or Hyp, C = Pro or Hyp, D = Gly, W = Z = Phe or Thi, X = Ser and Y = any aromatic amino acid of the D-configuration.

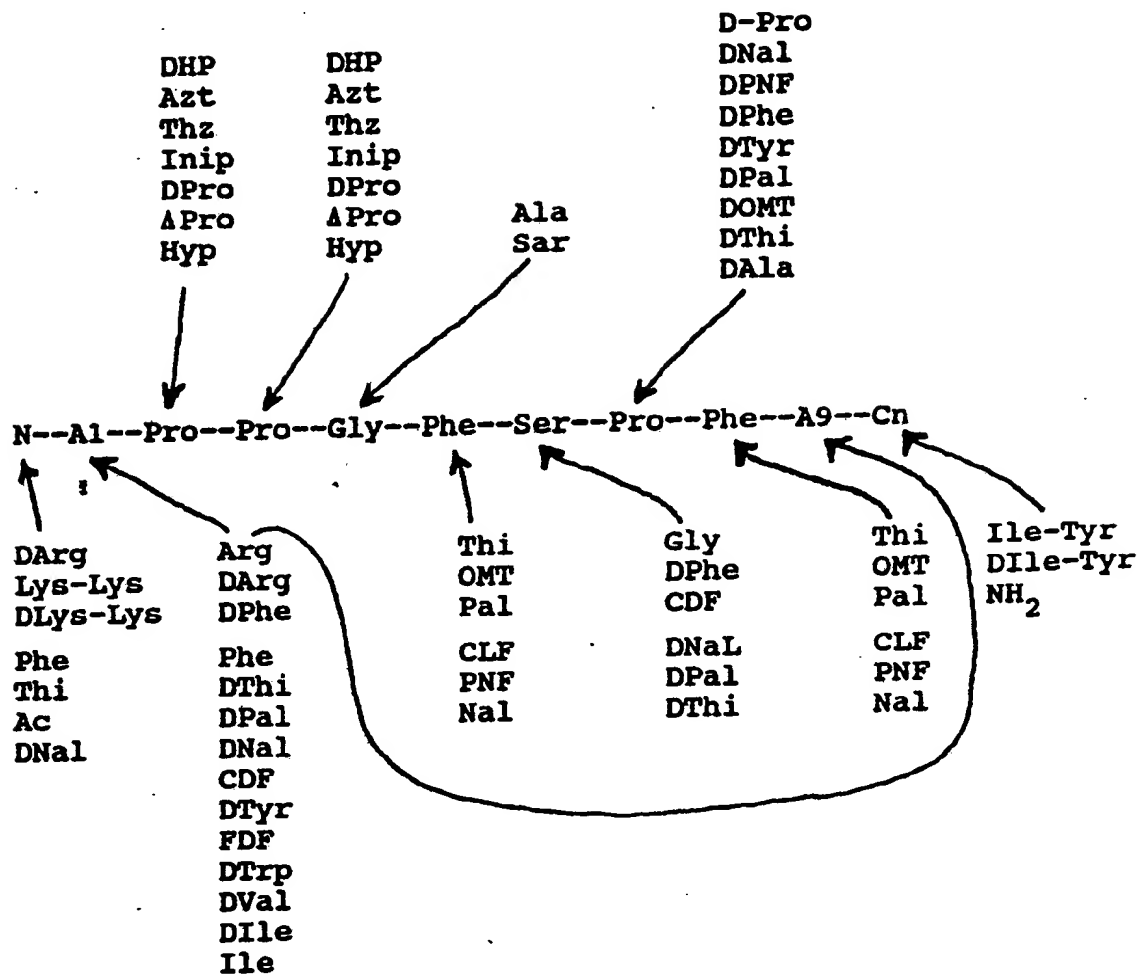
Salts of peptides of general formula I include salts with HCl, TFA, AcOH, as well as other pharmaceutically acceptable salts.

The following TABLES I and II show substitutions that can be made in the bradykinin polypeptide and the effect of such substitutions. Indicated substitutions of the 0, 1, 2, 3, 5, 7 and 8 amino acid residues of bradykinin yield preferred bradykinin antagonists.

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TABLE I

SUBSTITUTIONS IN BRADYKININ ANTAGONISTS



## TABLE I, Cont'd

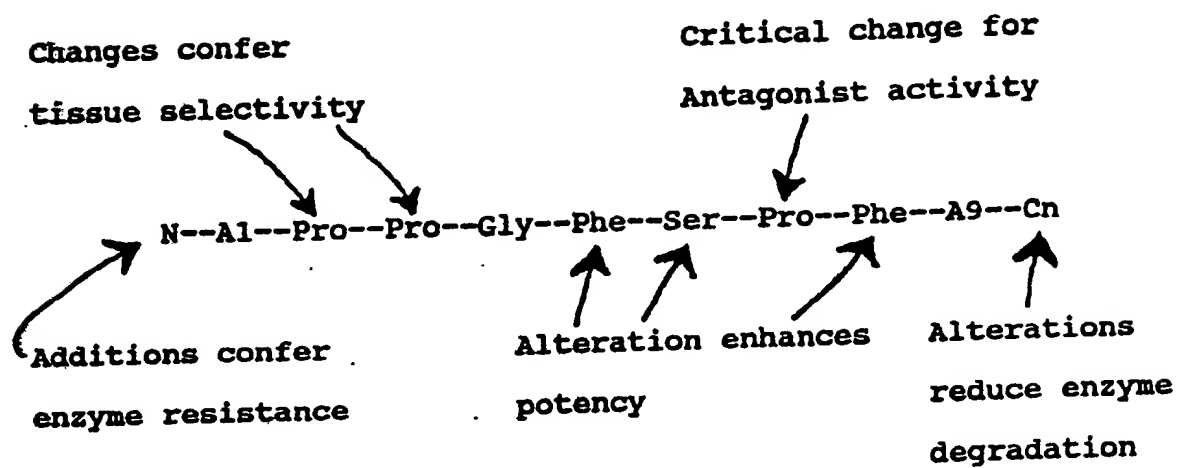
## SUBSTITUTIONS IN BRADYKININ ANTAGONISTS

Thi=  $\beta$ -(2-Thienyl)alanine  
Pal=  $\beta$ -(2-Pyridyl)alanine  
Hyp=4-Hydroxyproline  
Azt=Azetidine-2-carboxylic acid  
Thz=Thiazolidine-2-carboxylic acid  
Inip=Isonipecotic acid  
OMT=0-Methyltyrosine  
CDF=para-chloro-D-phenylalanine  
Nal=  $\beta$ -(2-Naphthyl)-alanine  
CLF=para-chloro-L-phenylalanine  
PNF=para-nitrophenylalanine  
 $\Delta$ Pro=2, 3-Dehydroproline  
DHP=3, 4-Dehydroproline

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TABLE II

## CHARACTERISTICS OF BRADYKININ ANTAGONISTS





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## DETAILED DESCRIPTION

The synthesis of the peptides of general Formula I, including derivation, activation, and coupling of protected amino acid residues, and their purification, and the analytical methods for determining identity and purity are included in the general body of knowledge of peptide chemistry, as described in Houben Weyl "Methoden der Organische Chemie" Vol. 16, parts I & II (1974) for solution-phase synthesis, and in "Solid Phase Peptide Synthesis" by Stewart and Young (1984) for synthesis by the solid-phase method of Merrifield. Any chemist skilled in the art of peptide synthesis can synthesize the peptides of general Formula I by standard solution methods or by manual or automated solid-phase methods.

The symbols and abbreviations used for amino acids, their derivatives and protecting groups, and peptides and their salts are those customarily used in peptide chemistry (Biochem. J. 126:773, 1972, the Journal reference is hereby incorporated by reference). For convenience several abbreviations are defined in Table III reproduced below. All amino acid residues, except Gly, described in the specification (but not the claims which claims cover compositions of the D- and L-configuration) are of the L-configuration unless otherwise specified.

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TABLE III

## ABBREVIATIONS FOR AMINO ACID RESIDUES

Aib	- alpha-aminoisobutyric acid
Ala	- alanine
Azt	- azetidine-2-carboxylic acid
CDF	- para-chloro-D-phenylalanine
CLF	- para-chloro-L-phenylalanine
DHP	- 3,4 dehydroproline
FDF	- para - phenylalanine-D phenylalanine
hPhe	- homo-phenylalanine
His	- histidine
Hyp	- 4-hydroxy-proline
Gly	- glycine
Ile	- Isoleucine
Inip	- isonipecotic acid
Leu	- leucine
MDY	- O-methyl-D-tyrosine
OMT	- O-methyl-tyrosine
Nal	- beta-(2-naphthyl)-alanine
APro	- 2,3-dehydroproline
Pal	- beta-(3-pyridyl)-alanine or Z pyredyl
Phg	- alpha-phenylglycine
Phe	- phenylalanine

TABLE III, Cont'd

## ABBREVIATIONS FOR AMINO ACID RESIDUES,

PNF	- para-nitrophenalanine
Sar	- sarcosine
Thi	- beta-(2-thienyl)-alanine
Thz	- thiazolidine-2-carboxylic acid
Tyr	- tyrosine

(all other abbreviations follow the IUPAC standards for amino acid residues)

The following examples are illustrative of compounds of this invention with general formula I and are not limitative. All percentages and ratios are by weight when solids are involved and by volume when only liquids are involved.

## EXAMPLE 1

Preparation of Arg-Pro-Pro-Gly-Phe-Ser-DPhe-Phe-Arg (DPhe<sup>7</sup>-BK).

A mixture of 6.4 gm of tertiary butyloxy carbonyl-(g-paratoluene sulfonyl)-Arg [Boc-Arg(Tos)](15mMole) and 183

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mg of N, N-dimethylaminopyridine (1.5mMole) was dissolved in a mixture of 20 ml of dimethylformamide (DMF) and 125 ml of dichloromethane (DCM). Fifteen g (grams) of hydroxymethyl-polystyrene- divinyl benzene (1% crosslinked, containing 0.74 mMole of free hydroxyl group per g of resin) was added, followed by 60 ml of a 0.25 M solution of dicyclohexylcarbodiimide (DCC) in DCM at room temperature. The suspension was stirred at room temperature overnight, filtered, and the resin was washed three times with 60 ml of DCM, three times with 60 ml of methyl alcohol (MeOH), and reswollen in 120 ml of DCM. The coupling of another portion of Boc-Arg(Tos) was conducted on the resin as above. After filtering and washing the resin it was reswollen in 120 ml of DCM, and 2.1 ml of benzoyl chloride and 1.5 ml of triethylamine ( $\text{Et}_3\text{N}$ ) were added. After stirring the suspension for 30 minutes at room temperature the resin was filtered, washed three times with 60 ml portions of DCM, MeOH, washed three times with 60 ml portion of MeOH and finally washed three times with 60 ml portions of DCM. The resin was air dried to constant weight to give 18.5 gm of Boc-Arg(Tos) -hydroxymethyl-resin, with an actual amino acid content of 0.272 millimoles of Arg per g of resin as determined by quantitative amino acid analysis of a sample of the amino acid resin following hydrolysis (4 hr, 130 degrees C) in 6 N HCL/pr pionic acid.

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The resin, 1.5 gm containing a total of 0.4 mMole of Arg, was placed in the reaction vessel of an automatic solid-phase synthesizer (Beckman model 990) and subjected to one cycle of addition for the coupling of Boc-Phe as follows;

**PROGRAM A. STANDARD DCC COUPLING:**

The resin was washed three times with 20ml portions of DCM. The resin was then equilibrated with 20ml of a 1:3 ratio of trifluoroacetic acid (TFA) in DCM containing 0.1% indole for 1.5 minutes. The equilibration was then repeated for 30 minutes. The resin was then washed six times with 20ml portions of DCM followed by neutralization with a 10% solution of (Et<sub>3</sub>N) in DCM for one and one half minutes, then the neutralization step was repeated. The resin was washed six times with 20ml of DCM and then equilibrated with a solution of 1.0 mMole of Boc-Phe in DCM for one and one half minutes. Then four ml of 0.25 N DCC in DCM was added and the mixture stirred for two hours. Then the resin was washed three times with 20ml portions of DCM.

A second cycle of addition was performed according to Program B:

**PROGRAM B. REVERSE ADDITION:**

The procedure of Program A through neutralization and following washes was repeated. Then 1.0 mMole of DCC in 4ml of

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DCM was added and the resin and solution were mixed for one and one-half minutes. Then 1.0 mMole of Boc-D-Phe in 12ml DCM was added and the resin and solution were mixed for two hours. The resin was then washed six times with 20ml portions of DCM.

The N-Terminal protecting group was removed according to the following sequence:

**PROGRAM C. TERMINAL DEPROTECTION:**

The procedure of PROGRAM A up to the neutralization with triethylamine was repeated. The resin was then washed 6 times with 20ml portions of ethyl alcohol and the peptide-resin was air dried giving 1.66 gm of DPhe-Phe-Arg-Resin as the trifluoroacetic acid salt.

Synthesis was continued with 410 mg of the DPhe-Phe-Arg-Resin TFA salt. The next residue was added according to PROGRAM D.

**PROGRAM D. RECOUPLE:**

The peptide-resin salt was first washed three times with 20ml portion of DCM, then neutralized with 10% Et<sub>3</sub>N DCM for 1.5 minutes. The neutralization step was then repeated and the peptide-resin-salt was washed six times with 20ml portions of DCM. The peptide-resin was then equilibrated with a solution of 1.0 mMole of Boc-Ser(OBzl) in DMF for 1.5 minutes.

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Four ml of 0.25 N DCC in DCM was added and mixed with the resin for two hours. The product was washed three times with DCM.

The following amino acid derivatives were added to the growing peptide chain according to the listed Programs: Boc-Phe (A), Boc-Gly (A), Boc-Pro (A), Boc-Pro (A), followed by recouple of Boc-Pro (D), Boc-Arg(Tos) (dissolved in 2 ml DMF + 9 ml DCM), (A), followed by Program C. This gave 530 mg of protected nonapeptide-resin as the TFA salt.

A 510 mg portion of the peptide-resin above was suspended in 10 ml of liquid anhydrous HF containing 1 ml of anisole at -70 degrees C and stirred 45 min. at 0 degrees C. HF and anisole were removed by vacuum (1 hr water pump, 1 hr vacuum pump), the peptide plus resin was washed three times with 20ml portions of ethyl ether ( $\text{Et}_2\text{O}$ ) and the peptide extracted into glacial acetic acid using three 6 ml extractions. The acetic acid solution was lyophilized to give 185 mg of crude deprotected peptide.

The peptide was purified by countercurrent distribution (CCD) (100 upper phase transfers in a Post CCD apparatus) in the solvent system nBuOH:1% TFA (1:1). The content of the tubes corresponding to the main peptide-containing peak, as determined by the quantitative Sakaguchi reagent, was collected, the solvent evaporated under reduced pressure, the residue dissolved in glacial acetic acid (AcOH)

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and lyophilized to give 140 mg of peptide with a partition coefficient (k) from the CCD of 5.7. Repeating the countercurrent distribution in the solvent system nBuOH:AcOH:H<sub>2</sub>O (4:1:5) gave, upon detection and workup as described above, 73 mg of Arg-Pro-Pro-Gly-Phe-Ser-DPhe-Phe-Arg as the TFA salt (k = 0.2). Thin layer chromatographs (TLC) on Merck glass precoated silica gel plates in the solvent systems nBuOH:AcOH:H<sub>2</sub>O (8:3:4) and EtOAc:pyridine:AcOH:H<sub>2</sub>O (5:5:1:3) gave R<sub>f</sub>(834) of 0.17 and R<sub>f</sub>(5513) of 0.36 for the pure peptide, as visualized by the chlorine-tolidine peptide identification spray. Quantitative amino acid analysis (Beckman 120 instrument) after acid hydrolysis (17 hr in sealed glass vials under N<sub>2</sub> at 110° C in 2 ml 6 N HCl containing 2 drops 2-mercaptoethanol and 40 microliters of phenol) gave the following ratios of amino acids: Arg(2.12); Pro(1.93); Gly(1.01); Phe(2.98); Ser(0.96).

#### EXAMPLES 2 - 40

Examples 2 - 40 represent bradykinin antagonists containing a hydrophobic amino acid residue at positions one and seven and were prepared by methods similar to those described in Example 1 and are not limitative.

2. CDF-Pro-Pro-Gly-Phe-Ser-DPhe-Phe-Arg (CDF<sup>1</sup>DPhe<sup>7</sup>-BK): k(415) = 2.57 Arg - 1.03, Pro - 1.95, Gly - 1.04, Phe - 2.98, Ser -



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0.94, PCF - 1.06.

3. CDF-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (CDF<sup>1</sup>Hyp<sup>3</sup>DPhe<sup>7</sup>-BK):  
k(415) = 1.947 Arg - 1.05, Pro - 1.00, Gly - 0.96, Phe - 3.02,  
Ser - 0.94, PCF - 1.10, Hyp - 0.99.

4. DArg-CDF-Pro-Pro-Gly-Phe-Ser-DPhe-Phe-Arg (DArg<sup>0</sup>CDF<sup>1</sup>-DPhe<sup>7</sup>-BK): k(415) = 0.754.

5. DArg-CDF-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg  
(DArg<sup>0</sup>CDF<sup>1</sup>Hyp<sup>3</sup>DPhe<sup>7</sup>-BK): k(415) = 1.272 Arg - 1.92, Pro - 1.03,  
Gly - 1.05, Phe - 3.04, Ser - 0.95, PCF - 1.00, Hyp - 1.00.

6. CDF-Pro-Pro-Gly-Phe-Ser-DThi-Phe-Arg (CDF<sup>1</sup>DThi<sup>7</sup>-BK): k(415)  
= 0.460 Arg - 1.08, Pro - 2.01, Gly - 1.03, Phe - 2.00, Ser -  
0.91, Thi - 0.93, PCF - 1.03.

7. DArg-CDF-Pro-Pro-Gly-Phe-Ser-DThi-Phe-Arg (DArg<sup>0</sup>CDF<sup>1</sup>DThi<sup>7</sup>-BK): k(415) = 0.754 Arg - 2.09, Pro - 1.92, Gly - 1.03,  
Phe - 2.10, Ser - 0.92, Thi - 0.91, PCF - 1.09.

8. DArg-Pro-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg (DArg<sup>1</sup>Hyp<sup>3</sup>Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK): k(1.1) = 2.279 Arg - 2.11, Pro - 1.04, Gly -  
1.03, Phe - 1.13, Ser - 0.75, Hyp - 1.55, Thi - 1.90.

9. DTyr-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (DTyr<sup>1</sup>Hyp<sup>3</sup>DPhe<sup>7</sup>-BK):  
k(415) = 1.326 Arg - 0.99, Pro - 1.06, Gly - 1.02, Phe - 3.02,  
Tyr - 1.01, Ser - 0.94, Hyp - 0.96.

10. DArg-DTyr-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (DArg<sup>0</sup>DTyr<sup>1</sup>Hyp<sup>3</sup>DPhe<sup>7</sup>-BK): k(415) = 0.408 Arg - 2.06, Pro - 1.06, Gly -  
1.04, Ph - 3.19, Ser - 0.88, Hyp - 0.92, Tyr - 0.85.

11. FDF-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (FDF<sup>1</sup>Hyp<sup>3</sup>DPhe<sup>7</sup>-BK):  
k(415) = 1.857 Arg - 1.00, Pro - 1.03, Gly - 1.05, Phe - 2.99,  
Ser - 0.93, FDF - 0.95, Hyp - 1.05.
12. DArg-FDF-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (DArg<sup>0</sup>FDF<sup>1</sup>  
Hyp<sup>3</sup>DPhe<sup>7</sup>-BK): k(415) = 0.538 Arg - 2.11, Pro - 1.02, Gly -  
1.05, Phe - 2.96, Ser - 0.88, Hyp - 0.94, FDF - 1.04.
13. DTrp-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (DTrp<sup>1</sup>Hyp<sup>3</sup>  
DPhe<sup>7</sup>-BK): k(415) = 8.091.
14. DArg-DTrp-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (DArg<sup>0</sup>DTrp<sup>1</sup>  
Hyp<sup>3</sup>DPhe<sup>7</sup>-BK): k(415) = 7.333.
15. DVal-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (DVal<sup>1</sup>Hyp<sup>3</sup>  
DPhe<sup>7</sup>-BK): k(415) = 0.754 Arg - 1.02, Pro - 0.95, Gly - 1.06,  
Phe - 3.18, Ser - 0.91, Hyp - 0.98, Val - 0.90.
16. DArg-DVal-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (DArg<sup>0</sup>DVal<sup>1</sup>  
Hyp<sup>3</sup>DPhe<sup>7</sup>-BK): k(415) = 0.370 Arg - 2.06, Pro - 0.94, Gly -  
0.99, Phe - 3.01, Ser - 0.95, Hyp - 1.02, Val - 1.01.
17. DIle-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (DIle<sup>1</sup>Hyp<sup>3</sup>  
DPhe<sup>7</sup>-BK): k(415) = 1.128 Arg - 1.01, Pro - 0.96, Gly - 0.97,  
Phe - 3.03, Ser - 0.98, Hyp - 1.03, Ile - 0.98.
18. DArg-DIle-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (DArg<sup>0</sup>DIle<sup>1</sup>  
Hyp<sup>3</sup>DPhe<sup>7</sup>-BK): k(415) = 0.639 Arg - 2.00, Pro - 0.92, Gly -  
1.01, Phe - 3.09, Ser - 0.98, Hyp - 1.01, Ile - 0.90.

## EXAMPLES 19 - 40

Examples 19 - 40 represent bradykinin antagonists containing hydrophobic amino acid residues at positions one and seven and were prepared by methods similar to those described in Example 1 and are not limitative.

19. DPhe-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg (DPhe<sup>1</sup>Thi<sup>5,8</sup>

DPhe<sup>7</sup>-BK): k(415) = 1.703 Arg - 1.05, Pro - 2.00, Gly - 1.04, Phe - 2.09, Thi - 1.91, Ser - 0.92.

20. DThi-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg (DThi<sup>1</sup>Thi<sup>5,8</sup>

DPhe<sup>7</sup>-BK): k(415) = 1.326 Arg - 1.08, Pro - 2.04, Gly - 1.02, Ser - 0.98, Phe - 1.01, Thi - 2.88.

21. DPal-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg (DPal<sup>1</sup>Thi<sup>5,8</sup>

DPhe<sup>7</sup>-BK): k(1.1) = 2.333 Arg - 1.06, Pro - 2.01, Gly - 1.05, Phe - 1.02, Ser - 0.96, Thi - 1.96, Pal - 0.99.

22. DNal-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg (DNal<sup>1</sup>Thi<sup>5,8</sup>

DPhe<sup>7</sup>-BK): k(415) = 3.167 Arg - 0.97, Pro - 2.09, Gly - 1.01, Phe - 1.02, Ser - 1.00, Thi - 1.89, Nal - 0.99.

23. DArg-DNal-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg (DArg<sup>0</sup>DNal<sup>1</sup>

Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK): k(415) = 1.273 Arg - 2.16, Pro - 2.01, Gly - 1.05, Phe - 1.01, Thi - 1.88, Nal - 0.95, Ser - 0.94.

24. Lys-Lys-DNal-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg (Lys-Lys-

DNal<sup>1</sup>Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK): k(415) = 0.190 Arg - 1.05, Lys -

2-13, Pro - 1.97, Phe - 0.98, Ser - 0.86, Gly - 1.01, Thi - 1.98, Nal - 1.00.

25. D-Nal-DNal-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg (DNal<sup>0,1</sup>Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK): k(415) = 7.333 Arg - 1.14, Pro - 1.90, Gly - 1.11, Phe - 1.08, Ser - 1.01, Thi - 1.93, Nal - 1.63.

26. DNal-Hyp-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg (DNal<sup>1</sup>Hyp<sup>2,3</sup>Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK): k(415) = 1.941 Arg - 1.10, Gly - 1.00, Phe - 0.99, Ser - 1.00, Thi - 1.94, Hyp - 1.89, Nal - 1.00.

27. DArg-DNal-Hyp-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg (DArg<sup>0</sup>DNal<sup>1</sup>Hyp<sup>2,3</sup>Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK): k(415) = 0.587 Arg - 1.99, Gly - 1.08, Ser - 0.99, Phe - 1.04, Thi - 1.99, Hyp - 1.89, Nal - 1.02.

28. Lys-Lys-DNal-Hyp-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg (Lys-Lys-DNal<sup>1</sup>Hyp<sup>2,3</sup>Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK): k(415) = 0.587 Arg - 1.99, Gly - 1.08, Ser - 0.99, Phe - 1.04, Thi - 1.99, Hyp - 1.89, Nal - 1.02.

29. DNal-Pro-Pro-Gly-Phe-Ser-DPhe-Phe-Arg (DNal<sup>1</sup>DPhe<sup>7</sup>-BK): k(415) = 3.000 Arg - 1.00, Nal - 0.86, Pro - 1.93, Gly - 1.14, Phe - 3.13, Ser - 0.95.

30. DNal-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (DNal<sup>1</sup>Hyp<sup>3</sup>DPhe<sup>7</sup>-BK): k(415) = 2.448 Arg - 1.07, Pro - 1.02, Gly - 1.07, Phe - 3.01, Ser - 0.95, Nal - 0.93.

31. DArg-DNal-Pro-Pro-Gly-Phe-Ser-DPhe-Phe-Arg (DArg<sup>0</sup>DNal<sup>1</sup>DPhe<sup>7</sup>-BK): k(415) = 1.128 Arg - 2.15, Pro - 1.95, Gly - 1.01, Phe - 3.01, Ser - 0.92, Nal - 0.96.

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32. DArg-DNal-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Arg (DArg<sup>0</sup>DNal<sup>1</sup>Hyp<sup>3</sup>DPhe<sup>7</sup>-BK): k(415) = 0.786 Arg - 2.08, Nal - 0.89, Pro - 0.97, Gly - 1.02, Phe - 3.16, Ser - 0.92, Hyp - 0.96.
33. DThi-Pro-Pro-Gly-Phe-Ser-DThi-Phe-Arg (DThi<sup>1,7</sup>-BK): k(415) = 1.70, Arg - 1.05, Pro - 2.07, Gly - 1.05, Phe - 2.05, Ser - 0.94, Thi - 1.84.
34. DThi-Pro-Hyp-Gly-Phe-Ser-DThi-Phe-Arg (DThi<sup>1,7</sup>Hyp<sup>3</sup>-BK): k(415) = 1.22 Arg - 0.97, Pro - 1.07, Gly - 0.95, Phe - 2.07, Ser - 0.98, Thi - 1.85, Hyp - 1.11.
35. DArg-DThi-Pro-Pro-Gly-Phe-Ser-DThi-Phe-Arg (DArg<sup>0</sup>DThi<sup>1,7</sup>-BK): k(415) = 0.449 Arg - 1.87, Pro - 1.87, Gly - 0.98, Phe - 2.19, Ser - 0.89, Thi - 1.89.
36. DArg-DThi-Pro-Hyp-Gly-Phe-Ser-DThi-Phe-Arg (DArg<sup>0</sup>DThi<sup>1,7</sup>Hyp<sup>3</sup>-BK): k(415) = 0.449 Arg - 1.99, Pro - 1.06, Gly - 1.07, Phe - 2.03, Ser - 0.95, Thi - 1.82, Hyp - 1.07.
37. DNal-Pro-Pro-Gly-Phe-Ser-DThi-Phe-Arg (DNal<sup>1</sup>DThi<sup>7</sup>-BK): k(415) = 3.00 Arg - 0.92, Pro - 2.08, Gly - 1.06, Phe - 2.05, Ser - 0.91, Thi - 1.00, Nal - 0.92.
38. DNal-Pro-Hyp-Gly-Phe-Ser-DThi-Phe-Arg (DNal<sup>1</sup>Hyp<sup>3</sup>DThi<sup>7</sup>-BK): k(415) = 2.448 Arg - 1.10, Pro - 1.05, Gly - 1.05, Phe - 2.13, Ser - 0.84, Hyp - 1.00, Thi - 0.8, Nal - 0.87.
39. DArg-DNal-Pro-Pro-Gly-Phe-Ser-DThi-Phe-Arg (DArg<sup>0</sup>DNal<sup>1</sup>

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DThi<sup>7</sup>-BK): k(415) = 0.887 Arg - 2.15, Nal - 0.96, Pro - 2.00, Gly - 1.02, Phe - 2.04, Ser - 0.93, Thi - 0.90.

40. DArg-DNal-Pro-Hyp-Gly-Phe-Ser-DThi-Phe-Arg (DArg<sup>0</sup>DNal<sup>1</sup>Hyp<sup>3</sup>DThi<sup>7</sup>-BK): k(415) = 1.272 Arg - 2.16, Nal - 1.02, Pro - 0.94, Gly - 1.06, Phe - 1.99, Ser - 0.91, Thi - 1.01.

## EXAMPLES 41 - 79

Examples 41 - 79 represent bradykinin antagonist peptides possessing C-terminal modifications and extensions in which arginine in position 9 has been replaced or in which an extension has been made to the C-terminal position. The compounds of Examples 41 - 79, unless otherwise noted were prepared by methods similar to those described in Example 1 and are not limitative.

41. Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-DArg (Thi<sup>5,8</sup>DPhe<sup>7</sup>DArg<sup>9</sup>-BK): k(415) = 0.266 Arg - 2.08, Pro - 2.00, Gly - 1.01, Phe - 1.01, Ser - 0.94, Thi - 1.96.

42. Ac-Arg-Pro-Pro-Gly-Thi-Ser-DPhe<sup>7</sup>-Thi-DArg (Ac-Thi<sup>5,8</sup>DPhe<sup>7</sup>DArg<sup>9</sup>-BK): k(415) = 0.754 Arg - 2.09, Pro - 2.03, Gly - 1.03, Phe - 0.96, Ser - 0.96, Thi - 1.93.

43. DArg-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-DArg (DArg<sup>0</sup>Thi<sup>5,8</sup>DPhe<sup>7</sup>DArg<sup>9</sup>-BK): k(1.1) = 2.226 Arg - 3.05, Pro - 2.01, Gly - 1.08, Phe - 0.94, Ser - 1.00, Thi - 1.93.

44. Lys-Lys-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-DArg (Lys-Lys-Thi<sup>5,8</sup>DPhe<sup>7</sup>DArg<sup>9</sup>-BK): k(1.1) = 0.282 Arg - 2.00, Lys - 2.00, Pro - 1.98, Gly - 1.07, Phe - 1.03, Ser - 1.01, Thi - 1.92.

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Examples 45 - 51 represent bradykinin antagonist peptides possessing a C-terminal phenylalanine (Phe). Peptide analogs possessing a C-terminal Phe residue are prepared by methods described in the Example 1 for the preparation of [DPhe<sup>7</sup>]-BK, except that the starting amino acid resin is Boc-Phe -hydroxymethyl-Resin (Boc-Phe-HMR), which is prepared similarly to Boc-Arg(Tos)-HMR above. Boc-Phe (1.325 g (5 mMole) and 61 mg (0.5 mMole) p-dimethylaminopyridine (DMAP) are added to 5.0 g hydroxymethyl-polystyrene-divinylbenzene (2% crosslinked) in 80 ml DCM. To this is added 5 ml of 1.0 M DCC in CHCl<sub>3</sub>, and the mixture is stirred 3.5 hr. The mixture is filtered and the amino acid resin is washed thoroughly with DCM, EtOH, and DCM, suspended in 80 ml DCM at 0 degrees, and treated with a mixture of 4.4 ml benzoyl chloride (37.5 mM) and 3.8 ml pyridine (47 mM) for 15 min., followed by 5 hr. stirring at room temperature. The resin is filtered, washed with DCM and air dried to give 5.5 g Boc-Phe-HMR, containing 0.485 mMole of Phe per gram of resin, as determined by quantitative amino acid analysis.

45. Arg-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Phe (Hyp<sup>3</sup>DPhe<sup>7</sup>Phe<sup>9</sup>-BK):

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$k(415) = 1.50$  Arg - 1.03, Pro - 0.96, Gly - 1.07, Phe - 3.96,  
Ser - 0.97, Hyp - 1.00.

46. DArg-Arg-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Phe (DArg<sup>0</sup>-Hyp<sup>3</sup>  
DPhe<sup>7</sup>Phe<sup>9</sup>-BK):  $k(415) = 0.429$  Arg - 2.01, Pro - 0.99, Gly -  
0.98, Phe - 3.89, Ser - 0.99, Hyp -.

47. DNal-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Phe (DNal<sup>1</sup>Hyp<sup>3</sup>DPhe<sup>7</sup>  
Phe<sup>9</sup>-BK):  $k(415) = 5.667$ .

48. DArg-DNal-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Phe (DArg<sup>0</sup>  
DNal<sup>1</sup>Hyp<sup>3</sup>DPhe<sup>7</sup>Phe<sup>9</sup>-BK):  $k(415) = 5.667$ .

49. DArg-Pro-Hyp<sup>3</sup>-Gly-Phe-Ser-DPhe-Phe-Phe (DArg<sup>1</sup>Hyp<sup>3</sup>  
DPhe<sup>7</sup>Phe<sup>9</sup>-BK):  $k(415) = 1.439$  Arg - 1.04, Pro - 1.00, Gly -  
1.07, Phe - 3.90, Ser - 0.94, Hyp - 1.04.

50. DArg-DArg-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Phe (DArg<sup>0,1</sup>  
Hyp<sup>3</sup>DPhe<sup>7</sup>Phe<sup>9</sup>-BK):  $k(415) = 0.493$  Arg - 2.07, Pro - 0.95, Gly -  
1.01, Phe - 4.02, Ser - 0.90, Hyp -.

The C-terminal peptide amides of Examples 51 - 58 were prepared employing the following procedures. In the reaction vessel of an automatic peptide synthesizer (Beckmann 990), or by manual solid phase methods, 5.0 g of commercial p-methyl-benzhydrylamine-polystyrene-divinylbenzene resin (MBHA-Resin), 1% crosslinked, is subjected to 3 X 1.5 min. washes with dichloromethane (DCM), 2 X 1.5 min. treatments with 10% Et<sub>3</sub>N in DCM and 6 X 1.5 min. washes with DCM. The resin is



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equilibrated with a solution of 2.15 g (5 mMole) of commercial Boc-Arg(Tos)-OH in DCM for 1.5 min., 20 ml of 0.25 M dicyclohexylcarbodiimide (DCC) in DCM is added and the mixture stirred at room temp. 2 hr. The amino acid-Resin is washed 6 X 1.5 min. with DCM, and stirred for 2 hr. with a mixture of 0.94 ml acetic anhydride and 1.40 ml Et<sub>3</sub>N in DMF. The mixture is filtered, washed 3 X 1.5 min. with DCM and air dried to give 6.68 g Boc-Arg(Tos)-MBHA-Resin containing 0.404 mMole of Arg per gram of resin, as determined by quantitative amino acid analysis on an aliquot.

The Boc-Arg(Tos)-MBHA-Resin is treated to cycles of amino acid addition with various protected amino acids by standard solid-phase methods as described in Example 1 to produce the peptide amide-Resins desired. The desired peptide amide is obtained following hydrogen fluoride cleavage of the peptide from the resin and purification as described in Example 1.

51. Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg-NH<sub>2</sub> [(Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK)-NH<sub>2</sub>]: k(415) = 0.149 Arg - 2.10, Pro - 1.91, Gly - 1.04, Phe - 1.04, Ser - 0.97, Thi - 1.94.

52. Ac-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg-NH<sub>2</sub> [(Ac-Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK)-NH<sub>2</sub>]: k(415) = 0.515 Arg - 1.97, Pro - 1.95, Gly - 1.00, Phe - 1.02, Ser - 0.97, Thi - 2.00.

53. DArg-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg-NH<sub>2</sub> [(DArg<sup>0</sup>

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Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK)-NH<sub>2</sub>]: k(1.1) = 1.222 Arg - 3.16, Pro - 1.94, Gly - 1.02, Phe - 1.00, Ser - 0.91, Thi - 1.97.

54. Lys-Lys-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg-NH<sub>2</sub>  
 [(Lys-Lys-Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK)-NH<sub>2</sub>]: k(1.1) = 0.220 Arg - 2.09, Lys - 2.11, Pro - 1.92, Gly - 1.01, Phe - 1.00, Ser - 0.92, Thi - 1.93.

55. Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-DArg-NH<sub>2</sub> [(Thi<sup>5,8</sup>DPhe<sup>7</sup>DArg<sup>9</sup>-BK)-NH<sub>2</sub>]: k(415) = 0.191 Arg - 2.05, Pro - 1.95, Gly - 1.04, Phe - 1.05, Ser - 0.95, Thi - 1.95.

56. Ac-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-DArg-NH<sub>2</sub> [(Ac-Thi<sup>5,8</sup>DPhe<sup>7</sup>DArg<sup>9</sup>-BK)-NH<sub>2</sub>]: k(415) = 0.567 Arg - 2.05, Pro - 2.04, Gly - 1.01, Phe - 1.00, Ser - 0.94, Thi - 1.97.

57. DArg-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-DArg-NH<sub>2</sub> [(DArg<sup>0</sup>Thi<sup>5,8</sup>DPhe<sup>7</sup>DArg<sup>9</sup>-BK)-NH<sub>2</sub>]: k(1.1) = 1.174 Arg - 3.19, Pro - 1.94, Gly - 1.01, Phe - 0.98, Ser - 0.93, Thi - 1.95.

58. Lys-Lys-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-DArg-NH<sub>2</sub>  
 [(Lys-Lys-Thi<sup>5,8</sup>DPhe<sup>7</sup>DArg<sup>9</sup>-BK)-NH<sub>2</sub>]: k(1.1) = 0.205 Arg - 1.95, Lys - 2.00, Pro - 2.00, Gly - 1.04, Phe - 1.01, Ser - 1.05, Thi - 1.95.

Examples 59 - 78 represent bradykinin antagonist peptides possessing a C-terminal amino acid and peptide extension.

59. Arg-Pro-Pro-Gly-Phe-Ser-Pro-Phe-Arg-Ile-Tyr [(BK)-Ile

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-Tyr]:  $k(415) = 0.493$  Arg - 1.82, Pro - 3.13, Gly - 1.15, Phe - 1.99, Ser - 1.05, Ile - 0.86, Tyr - 1.00.

60. DArg-Pro-Pro-Gly-Phe-Ser-Pro-Phe-Arg-Ile-Tyr [(DArg<sup>0</sup>-BK)-Ile-Tyr]:  $k(415) = 0.176$  Arg - 2.87, Pro - 3.06, Gly - 1.12, Phe - 1.97, Ser - 1.08, Ile - 2.84, Tyr - 0.90.

61. Lys-Lys-Arg-Pro-Pro-Gly-Phe-Ser-Pro-Phe-Arg-Ile-Tyr [(Lys-Lys-BK)-Ile-Tyr]:  $k(1.1) = 1.439$  Arg - 1.96, Lys - 2.24, Pro - 3.01, Phe - 2.04, Gly - 1.01, Ser - 0.96, Ile - 0.85, Tyr - 0.93.

62. DLys-Lys-Arg-Pro-Pro-Gly-Phe-Ser-Pro-Phe-Arg-Ile-Tyr [(DLys-Lys-BK)-Ile-Tyr]:  $k(1.1) = 1.439$  Arg - 2.04, Lys - 2.05, Pro - 3.06, Phe - 2.08, Ser - 0.92, Gly - 1.05, Ile - 0.84, Tyr - 0.95.

63. Arg-Pro-Pro-Gly-Phe-Ser-DPhe-Phe-Arg-Ile-Tyr [(DPhe<sup>7</sup>-BK)-Ile-Tyr]:  $k(415) = 0.639$  Arg - 2.07, Pro - 2.00, Gly - 1.04, Phe - 3.05, Ser - 0.93, Ile - 0.93, Tyr - 0.98.

64. DArg-Pro-Pro-Gly-Phe-Ser-DPhe-Phe-Arg-Ile-Tyr [(DArg<sup>0</sup> DPhe<sup>7</sup>-BK)-Ile-Tyr]:  $k(415) = 0.220$  Arg - 2.80, Pro - 2.09, Gly - 1.17, Phe - 3.15, Ser - 0.96, Ile - 0.88, Tyr - 0.95.

65. Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg-Ile-Tyr [(Thi<sup>5,8</sup> DPhe<sup>7</sup>-BK)-Ile-Tyr]:  $k(415) = 0.613$  Arg - 2.06, Pro - 2.04, Gly - 1.12, Phe - 1.00, Ser - 0.95, Thi - 1.99, Ile - 0.90, Tyr - 0.94.

66. DArg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg-Ile-Tyr [(DArg<sup>0</sup>

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Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK)-Ile-Tyr]: k(415) = 0.235 Arg - 3.10, Pro - 1.98, Gly - 1.10, Thi - 1.87, Ser - 1.00, Ile - 0.88, Tyr - 1.05.

67. Arg-Pro-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg-Ile-Tyr [(Hyp<sup>3</sup>Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK)-Ile-Tyr]: k(415) = 0.493 Arg - 2.05, Pro - 1.12, Gly - 1.08, Phe - 0, Thi - 1.86, Ser - 1.05, Hyp - 0.99, Ile - 0.84, Tyr - 0.97.

68. DArg-Pro-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg-Ile-Tyr [(DArg<sup>0</sup>Hyp<sup>3</sup>Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK)-Ile-Tyr]: k(415) = 13.236 Arg - 3.07, Pro - 1.05, Gly - 1.07, Phe - 0.99, Ser - 0.96, Thi - 1.96, Hyp - 0.98, Ile - 0.96, Tyr - 0.97.

69. Arg-Pro-Pro-Gly-Phe-Ser-DPhe-Phe-Arg-DIle-Tyr [(DPhe<sup>7</sup>-BK)-DIle-Tyr]: k(415) = 0.639 Arg - 2.12, Pro - 2.13, Gly - 0.94, Phe - 3.16, Ser - 1.00, Ile - 0.77, Tyr - 0.88.

70. DArg-Pro-Pro-Gly-Phe-Ser-DPhe-Phe-Arg-DIle-Tyr [(DArg<sup>0</sup>DPhe<sup>7</sup>-BK)-DIle-Tyr]: k(415) = 0.20 Arg - 2.86, Pro - 1.87, Gly - 1.01, Phe - 3.42, Ser - 0.71, Ile - 0.60, Tyr - 1.07.

71. Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg-DIle-Tyr [(Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK)-DIle-Tyr]: k(415) = 0.515 Arg - 2.03, Pro - 1.95, Gly - 1.07, Phe - 1.07, Ser - 0.88, Thi - 1.91, Ile - 0.84, Tyr - 1.09.

72. DArg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg-DIle-Tyr [(DArg<sup>0</sup>Thi<sup>5,8</sup>DPhe<sup>7</sup>-BK)-DIle-Tyr]: k(415) = 0.205 Arg - 3.15, Pro - 2.17, Gly - 0.99, Phe - 1.02, Ser - 1.00, Thi - 1.86, Ile -

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- 0.83, Tyr - 1.02.

73. Arg-Pro-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg-DIle-Tyr [(Hyp<sup>3</sup> Thi<sup>5,8</sup> DPhe<sup>7</sup>-BK)-DIle-Tyr]: k(415) = 0.429 Arg - 2.06, Pro - 1.13, Gly - 1.01, Phe 1.07, Ser - 0.90, Thi - 2.00, Ile - 0.89, Tyr - 1.03.

74. DArg-Pro-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg-DIle-Tyr [(DArg<sup>0</sup> Hyp<sup>3</sup> Thi<sup>5,8</sup> DPhe<sup>7</sup>-BK)-DIle-Tyr]: k(415) = 0.735 Arg - 3.28, Pro - 1.11, Gly - 1.10, Phe - 1.07, Ser - 0.90, Hyp - 0.80, Thi - 1.78, Ile - 0.85, Tyr - 1.11.

75. DNal-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg-DIle-Tyr [(DNal<sup>1</sup> Thi<sup>5,8</sup> DPhe<sup>7</sup>-BK)-DIle-Tyr]: k(415) = 10.111 Arg - 1.21, Pro - 1.75, Gly - 1.13, Phe - 1.19, Ser - 0.97, Thi- 1.79, Nal - 0.75, Ile - 0.90, Tyr - 1.3.

76. DArg-DNal-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg-DIle-Tyr [(DArg<sup>0</sup> DNal<sup>1</sup> Thi<sup>5,8</sup> DPhe<sup>7</sup>-BK)-DIle-Tyr]: k(415) = 2.333 Arg - 2.07, Pro - 2.07, Gly - 1.04, Phe - 1.05, Ser - 0.86, Tyr - 0.96, Ile - 1.07, Nal - 1.02, Thi - 1.99.

77. DNal-Pro-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg-DIle-Tyr [(DNal<sup>1</sup> Hyp<sup>3</sup> Thi<sup>5,8</sup> DPhe<sup>7</sup>-BK)-DIle-Tyr]: k = Arg - 1.04, Pro - 1.01, Phe - 1.15, Ser - 0.98, Hyp - 0.98, Nal - 0.93, Thi - 1.84, Ile - 0.98, Tyr - 1.06.

78. DArg-DNal-Pro-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg-DIle-Tyr [(DArg<sup>0</sup> DNal<sup>1</sup> Hyp<sup>3</sup> Thi<sup>5,8</sup> DPhe<sup>7</sup>-BK)-DIle-Tyr]: k(415) = Arg - 2.11, Pro - 0.95, Gly - 1.00, Phe - 1.08, Ser - 0.99, Hyp -

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1.01, Thi - 1.93, Tyr - 0.94, Ile - 0.95.

#### EXAMPLES OF BRADYKININ ANTAGONIST ACTIVITY

The bradykinin antagonists were assayed on isolated rat uterus in natural or induced estrus and on guinea pig ileum, according to the commonly accepted assay methods for bradykinin and related kinins as described by Trautschold (Handbook of Expt. Pharmacol. Vol 25, Springer Verlag, pp. 53-55, 1970) for inhibition of the myotropic activity of bradykinin. The inhibition potencies, as determined according to the commonly accepted manner described by Schild for antagonists of biologically active compounds (Br. J. Pharmacol. 2:189, 1947), and expressed as  $PA_2$  values are determined on isolated rat uterus (RUT) and isolated guinea pig ileum (GPI). In the assays, a dose-response curve is determined for the reference substance bradykinin. The dose of bradykinin which produced a half maximal contraction of tissue is the  $ED_{50}$  dose. An amount of bradykinin equivalent to twice the  $ED_{50}$  dose is administered to the tissue 30 seconds after the start of incubation of the tissue with a dose of antagonist. Doses of antagonist are increased in this protocol until pre-incubation with a dose of antagonist reduces the contraction in response to the double  $ED_{50}$  dose of bradykinin to response of a single  $ED_{50}$  dose of bradykinin. The  $PA_2$  value represents the negative

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logarithm of the molar concentration of antagonist necessary to reduce the response of a double  $ED_{50}$  dose of bradykinin to that of an  $ED_{50}$  dose. One unit of  $pa_2$  value represents an order of magnitude change in potency. For comparison, the negative log of the dose of BK, the dose which causes half maximal contraction of the tissues, is commonly known as the  $pd_2$  value. The  $pd_2$  value for bradykinin is 7.9 on the rat uterus and 7.4 on the guinea pig ileum. The values for compounds of various Examples are reported in Table IV.

TABLE IV  
BRADYKININ ANTAGONISM ON SMOOTH MUSCLES

EXAMPLE No.	STRUCTURE	$pa_2$ /RUT	$pa_2$ /GPI
1.	DPhe <sup>7</sup> -BK		<u>5.0</u>
2.	(CDF <sup>1</sup> DPhe <sup>7</sup> )-BK	<u>5.0(6)</u>	0
3.	(CDF <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	MIXED	0
4.	DArg-(CDF <sup>1</sup> DPhe <sup>7</sup> )-BK	<u>5.4(8)</u>	0
5.	DArg-(CDF <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	<u>5.8(8)</u>	0
6.	(CDF <sup>1</sup> DThi <sup>7</sup> )-BK	<u>I(2)</u>	0
7.	DArg-(CDF <sup>1</sup> DThi <sup>7</sup> )-BK	0	<u>I(P)</u>
8.	(DArg <sup>1</sup> Hyp <sup>3</sup> Thi <sup>5,8</sup> -DPhe <sup>7</sup> )-BK	<u>5.4(6)</u>	<u>5.4(2)</u>
9.	(DTyr <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	I	0
10.	DArg-(DTyr <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK		I
11.	(FDF <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	I	0

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TABLE IV, Cont'd

## BRADYKININ ANTAGONISM ON SMOOTH MUSCLES

EXAMPLE NO.	STRUCTURE	PA <sub>2</sub> /RUT	PA <sub>2</sub> /GPI
12.	DArg-(FDF <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	<u>5.7(4)</u>	0
13.	(DTrp <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK		
14.	DArg-(DTrp <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK		
15.	(DVal <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	0	0
16.	DArg-(DVal <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	<u>I(P)</u>	0
17.	(DIle <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	I	0
18.	DArg-(DIle <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	I	0
19.	(DPhe <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	0	0
20.	(DThi <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	0	0
21.	(DPal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	0	0
22.	(DNal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-Bk	<u>5.6(8)</u>	0
23.	DArg-(DNal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	<u>5.8(6)</u>	0
24.	Lys-Lys-(DNal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	<u>I(2X)</u>	0
25.	DNal-(DNal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	<u>5.6(6)</u>	
26.	(DNal <sup>1</sup> Hyp <sup>2,3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-Bk	0	0
27.	DArg-(DNal <sup>1</sup> Hyp <sup>2,3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	<u>5.3(6)</u>	0
28.	Lys-Lys-(DNal <sup>1</sup> Hyp <sup>2,3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	0	0
29.	(DNal <sup>1</sup> DPh <sup>7</sup> )-Bk	<u>5.4(6)</u>	0
30.	(DNal <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-Bk	0	0.1±



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TABLE IV, Cont'd

## BRADYKININ ANTAGONISM ON SMOOTH MUSCLES

EXAMPLE NO.	STRUCTURE	pA <sub>2</sub> /RUT	pA <sub>2</sub> /GPI
31.	DArg-(DNal <sup>1</sup> DPhe <sup>7</sup> )-BK	5.6(5)	0
32.	DArg-(DNal <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	5.9(6)	0
33.	(DThi <sup>1</sup> DThi <sup>7</sup> )-BK	0	0
34.	(DThi <sup>1</sup> Hyp <sup>3</sup> DThi <sup>7</sup> )-BK	0	I/0
35.	DArg-(DThi <sup>1</sup> DThi <sup>7</sup> )-BK	0	0
36.	DArg-(DThi <sup>1</sup> Hyp <sup>3</sup> DThi <sup>7</sup> )-BK	I/0	0
37.	(DNal <sup>1</sup> DThi <sup>7</sup> )-BK	0	0
38.	(DNal <sup>1</sup> Hyp <sup>3</sup> DThi <sup>7</sup> )-BK	0	0
39.	DArg-(DNal <sup>1</sup> DThi <sup>7</sup> )-BK	<u>5.6(4)</u>	0
40.	DArg-(DNal <sup>1</sup> Hyp <sup>3</sup> DThi <sup>7</sup> )-BK	<u>5.6(4)</u>	0
41.	(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK	0.01%	0
42.	Ac-(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK	0	0
43.	DArg-(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK	0	0
44.	Lys-Lys-(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK	0	0
45.	(Hyp <sup>3</sup> DPhe <sup>7</sup> Phe <sup>9</sup> )-BK		
46.	DArg-(Hyp <sup>3</sup> DPhe <sup>7</sup> Phe <sup>9</sup> )-BK		
47.	(DNal <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> Phe <sup>9</sup> )-BK		
48.	DArg-(DNal <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> Phe <sup>9</sup> )-BK		
49.	(DArg <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> Phe <sup>9</sup> )-BK		
50.	DArg-(DArg <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> Phe <sup>9</sup> )-BK		

TABLE IV, Cont'd

## BRADYKININ ANTAGONISM ON SMOOTH MUSCLES

EXAMPLE NO.	STRUCTURE	PA <sub>2</sub> /RUT	PA <sub>2</sub> /GPI
51.	(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-NH <sub>2</sub>	0.02%	0
52.	Ac-(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-NH <sub>2</sub>	0	0
53.	DArg-(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-NH <sub>2</sub>	0	I/0
54.	Lys-Lys-(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-NH <sub>2</sub>	0.02%	0
55.	(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK-NH <sub>2</sub>	0.01%	0
56.	Ac(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK-NH <sub>2</sub>	0	0
57.	DArg-(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK-NH <sub>2</sub>	0.003%	0
58.	Lys-Lys(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK-NH <sub>2</sub>	0.02%	0
59.	BK-Ile-Tyr	26%	21%
60.	DArg-BK-Ile-Tyr	25%	34%
61.	Lys-Lys-BK-Ile-Tyr	18%	21%
62.	DLys-Lys-BK-Ile-Tyr	5%	1%
63.	(DPhe <sup>7</sup> )-BK-Ile-Tyr	0.6%	MIXED
64.	DArg-(DPhe <sup>7</sup> )-BK-Ile-Tyr	0.7%	<u>6.2(5)</u>
65.	(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-Ile-Tyr	0.5%	MIXED
66.	DArg-(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-Ile-Tyr	0.3%	<u>5.8(5)</u>
67.	(Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-Ile-Tyr	0.06%	0

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TABLE IV, Cont'd

## BRADYKININ ANTAGONISM ON SMOOTH MUSCLES

EXAMPLE NO.	STRUCTURE	pA <sub>2</sub> /RUT	pA <sub>2</sub> /GPI
68.	DArg-(Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-Ile-Tyr	0.02%	6.7(5)
69.	(DPhe <sup>7</sup> )-BK-DIle-Tyr	0	0
70.	DArg-(DPhe <sup>7</sup> )-BK-DIle-Tyr	I/0	0
71.	(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr	0	0
72.	DArg-(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr	0	0
73.	(Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr	0.04%	0
74.	DArg-(Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr	0	0
75.	(DNaI <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr	I	I
76.	DArg-(DNaI <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr		0
77.	(DNaI <sup>1</sup> Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr		0
78.	DArg-(DNaI <sup>1</sup> Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr		0

Biological activity is listed for the analogs on rat uterus (RUT), guinea pig ileum (GPI). Agonist potency is

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Listed as percent of BK potency. Antagonist potency is listed as the  $PA_2$  value and is underlined, followed in parentheses by the number of tissues in the determination. I(#) indicates antagonism in # determinations without quantitation. I/0 indicates analog exhibits both antagonism and no effect on separate tissues in screening assays. MIXED indicates a mixed agonist/antagonist analog.

#### EXAMPLE OF THE ANTAGONISM OF BRADYKININ ANTAGONISTS ON RAT BLOOD PRESSURE

The in vivo effects of bradykinin antagonists on blood pressure in the anesthetized rat are determined according to the assay described by Roblero, Ryan and Stewart (Res. Commun. Pathol. Pharmacol. 6:207, 1973). The antagonists also produce inhibition of the bradykinin response when injected as a bolus admixture of bradykinin plus antagonist by either the ia or iv route of administration. The results of tests on compounds of the various Examples are - reported in Table V.

TABLE V

## BRADYKININ ANTAGONISM ON BLOOD PRESSURE

EXAMPLE NO	STRUCTURE	RBP-IA	RBP-IV	% DESTR
1.	DPhe <sup>7</sup> -BK			
2.	(CDF <sup>1</sup> DPhe <sup>7</sup> )-BK	<u>I(B)</u>	<u>I(B)</u>	-
3.	(CDF <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	<u>I(P)</u>	<u>I(P)</u>	-
4.	DArg-(CDF <sup>1</sup> DPhe <sup>7</sup> )-BK	<u>I(P)</u>	<u>I(P)</u>	-
5.	DArg-(CDF <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	<u>I(P)</u>	<u>I(P)</u>	-
6.	(CDF <sup>1</sup> DThi <sup>7</sup> )-BK	0	0	-

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TABLE V, Cont'd

## BRADYKININ ANTAGONISM ON BLOOD PRESSURE

EXAMPLE NO.	STRUCTURE	RBP/IA	RBP-IV	%DEST
7.	DArg-(CDF <sup>1</sup> DThi <sup>7</sup> )-BK	0	0	-
8.	(DArg <sup>1</sup> Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	<u>I(B)</u>	<u>I(B)</u>	-
9.	(DTyr <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	<u>I(B)</u>	<u>I(B)</u>	-
10.	DArg-(DTyr <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	<u>I(B)</u>	<u>I(B)</u>	-
11.	(FDF <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	<u>I(B)</u>	<u>I(B)</u>	-
12.	DArg-(FDF <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	<u>I(B)</u>	<u>I(B)</u>	-
13.	(DTrp <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK			
14.	DArg-(DTrp <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK			
15.	(DVal <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	0	0	-
16.	DArg-(DVal <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	0	0	-
17.	(DIle <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	0	0	-
18.	DArg-(DIle <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	0	0	-
19.	(DPhe <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	<u>I(B)</u>	<u>I(B)</u>	-
20.	(DThi <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	0	0	-
21.	(DPal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	0	0	-
22.	(DNal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	<u>I(B)</u>	<u>I(B)</u>	-
23.	DArg-(DNal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	<u>I(B)</u>	<u>I(B)</u>	-
24.	Lys-Lys-(DNal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	<u>I(B)</u>	<u>I(B)</u>	-
25.	DNal-(DNal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK			
26.	(DNal <sup>1</sup> Hyp <sup>2,3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	<u>I(B)</u>	<u>I(B)</u>	-

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TABLE V, Cont'd

## BRADYKININ ANTAGONISM ON BLOOD PRESSURE

EXAMPLE NO.	STRUCTURE	RBP-IA	RBP-IV	% DESTR
27.	DArg-(DNaI <sup>1</sup> Hyp <sup>2,3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	I(B)	I(B)	-
28.	Lys-Lys-(DNaI <sup>1</sup> Hyp <sup>2,3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK	I(B)	I(B)	-
29.	(DNaI <sup>1</sup> DPhe <sup>7</sup> )-BK	I(B)	I(B)	-
30.	(DNaI <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	0	0	-
31.	DArg-(DNaI <sup>1</sup> DPhe <sup>7</sup> )-BK	I(B)	I(B)	-
32.	DArg-(DNaI <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> )-BK	I(B)	I(B)	-
33.	(DThi <sup>1</sup> DThi <sup>7</sup> )-BK	0	0	-
34.	(DThi <sup>1</sup> Hyp <sup>3</sup> DThi <sup>7</sup> )-BK	I(B)	I(B)	-
35.	DArg-(DThi <sup>1</sup> DThi <sup>7</sup> )-BK	0	0	-
36.	DArg-(DThi <sup>1</sup> Hyp <sup>3</sup> DThi <sup>7</sup> )-BK	I(B)	I(B)	-
37.	(DNaI <sup>1</sup> DThi <sup>7</sup> )-BK	I(B)	I(B)	-
38.	(DNaI <sup>1</sup> Hyp <sup>3</sup> DThi <sup>7</sup> )-BK	I(B)	I(B)	-
39.	DArg-(DNaI <sup>1</sup> DThi <sup>7</sup> )-BK	0	0	-
40.	DArg-(DNaI <sup>1</sup> Hyp <sup>3</sup> DThi <sup>7</sup> )-BK	I(B)	I(B)	-
41.	(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK	0.1%	0.6%	77%
42.	Ac-(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK	0	0	-
43.	DArg-(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK	I(B)	I(B)	-
44.	Lys-Lys-(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK	0.2%	14%	0
45.	(Hyp <sup>3</sup> DPhe <sup>7</sup> Phe <sup>9</sup> )-BK	0	0	-

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TABLE V, Cont'd

## BRADYKININ ANTAGONISM ON BLOOD PRESSURE

EXAMPLE NO.	STRUCTURE	RBP-IA	RBP-IV	% DESTR
46.	DArg-(Hyp <sup>3</sup> DPhe <sup>7</sup> Phe <sup>9</sup> )-BK	I(B)	I(B)	-
47.	(DNaI <sup>1</sup> -Hyp <sup>3</sup> -DPhe <sup>7</sup> -Phe <sup>9</sup> )-BK			
48.	DArg-(DNaI <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> Phe <sup>9</sup> )-BK			
49.	(DArg <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> Phe <sup>9</sup> )-BK	0	0	-
50.	DArg-(DArg <sup>1</sup> Hyp <sup>3</sup> DPhe <sup>7</sup> Phe <sup>9</sup> )-BK	I(B)	I(B)	-
51.	(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-NH <sub>2</sub>	0.05%	2%	52%
52.	Ac-(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-NH <sub>2</sub>	I(B)	I(B)	-
53.	DArg-(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-NH <sub>2</sub>	0	I(P)	-
54.	Lys-Lys-(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-NH <sub>2</sub>	(long duration depressor)		
55.	(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK-NH <sub>2</sub>	0.1%	0.9%	61%
56.	Ac(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK-NH <sub>2</sub>	I(B)	I(B)	-
57.	DArg-(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK-NH <sub>2</sub>	I(B)	I(B)	-
58.	Lys-Lys(Thi <sup>5,8</sup> DPhe <sup>7</sup> DArg <sup>9</sup> )-BK-NH <sub>2</sub>	0.2%	3%	60%
59.	BK-Ile-Tyr	6%	103%	61%
60.	DArg-BK-Ile-Tyr	12%	189%	47%
61.	Lys-Lys-BK-Ile-Tyr	9%	497%	0
62.	DLys-Lys-BK-Ile-Tyr	5%	165%	0
63.	(DPhe <sup>7</sup> )-BK-Ile-Tyr	0.2%	2%	70%

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TABLE V, Cont'd

## BRADYKININ ANTAGONISM ON BLOOD PRESSURE

EXAMPLE NO.	STRUCTURE	RBP-IA	RBP-IV	% DESTR
64.	DArg-(DPhe <sup>7</sup> )-BK-Ile-Tyr	0	0	-
65.	(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-Ile-Tyr	0.06%	6%	0
66.	DArg-(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-Ile-Tyr	<u>I(B)</u>	<u>I(B)</u>	-
67.	(Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-Ile-Tyr	0	0	-
68.	DArg-(Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-Ile-Tyr	7%	AG	-
69.	(DPhe <sup>7</sup> )-BK-DIle-Tyr	0.3%	2%	75%
70.	DArg-(DPhe <sup>7</sup> )-BK-DIle-Tyr	0	0	-
71.	(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr	0.2%	2%	55%
72.	DArg-(Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr	0	0	-
73.	(Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr	0	0	-
74.	DArg-(Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr	<u>I(B)</u>	<u>I(B)</u>	-
75.	(DNal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr			
76.	DArg-(DNal <sup>1</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr	0	0	-
77.	(DNal <sup>1</sup> Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr			
78.	DArg-(DNal <sup>1</sup> Hyp <sup>3</sup> Thi <sup>5,8</sup> DPhe <sup>7</sup> )-BK-DIle-Tyr	<u>I(P)</u>	<u>I(P)</u>	-



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Biological activity is listed for the analogs on rat blood pressure (RBP) following intra-aortic (IA) and intravenous (IV) bolus administration. % DESTR indicates metabolic breakdown of BK-like agonists in RBP assay relative to BK = 100%. Agonist potency is listed as percent of BK potency. I(B) indicates antagonism of BK-induced depressor effect. MIXED indicates a mixed agonist/antagonist analog. AG indicates agonist activity. I(P) indicates BK antagonist activity as well as pressor activity.

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Therapeutic applications of the novel bradykinin antagonists include not only treatment for the production of bradykinin or related kinins by the animal but also the injection of bradykinin related peptides into an animal as a result of bites and stings or injection similarly of enzymes which cause production of bradykinin. Topical application alone or in combination with subcutaneous utilization of the bradykinin antagonists of the invention can be employed to treat the effects of bradykinin-related peptides causing pain, inflammation and swelling.

The therapeutic use of bradykinin antagonists of this invention for other traumatic, inflammatory or pathological conditions which are known to be mediated by bradykinin or exacerbated by an overproduction of bradykinin can also be achieved. These conditions include local trauma

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such as wounds, burns and rashes, angina, arthritis, asthma, allergies, rhinitis, shock, inflammatory bowel disease, low blood pressure, systemic treatment of pain and inflammation. The present bradykinin antagonists, as discussed, may be advantageously administered in a variety of ways including sublingual absorption as with nitroglycerine or patch administration using agents for assisting absorption through the skin such as for the treatment of angina. Based upon the  $PA_2$  and  $ED_{50}$  data disclosed in this invention and in the prior art related to agonist potency, it is possible for one skilled in the art to make a determination of the dosage of the novel bradykinin antagonists of the invention.

It is therefore estimated that the dosage range for typical application in such conditions as the pain and inflammation of wounds, burns and rashes would be 0.1 - 5 mg/ml; for a nasal spray formulation suitable for treating rhinitis, allergies and asthma suitable dosage range would be 0.1 - 5 mg/ml; for intravenous formulation suitable for the treatment of systemic inflammation, shock, arthritis, allergies, asthma; for an oral formulation for the treatment of inflammatory bowel disease or general pain and inflammation a suitable dosage range would be 10-100 mg/kg. Bradykinin

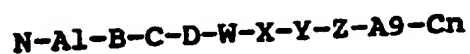
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antagonists can also be administered intravaginally, intrarectally, intrabuccally or by any other accepted internal application.

As will be recognized by those skilled in the art the present invention has a wide range of applicability to providing competitive inhibitors to the biological activities of bradykinin produced by the body in illness, injury and shock. The advantages of the invention in substituting the L-Pro position 7 with amino acids of the D-configuration to convert bradykinin agonists to antagonists provide a wide variety of specific and competitive antagonists for reducing the known effects of bradykinin. The additional advantages of the invention of modifying the L-Pro position 7 in conjunction with modifications at the other positions of the novel bradykinin antagonists provides a variety of useful compounds. It will further be appreciated that the present invention is susceptible to these and other modifications within the parameters of the invention without departing from the scope of the following claims.

## WHAT IS CLAIMED IS:

1                   1. A modified bradykinin type peptide of the  
2 formula:



3  
4 and the pharmaceutically acceptable salts thereof wherein

5                   (a) N is hydrogen, an amino acid residue of the D-  
6 or L- configuration, an N-terminal acyl type protecting group,  
7 an N-terminal aromatic or aliphatic urethane protecting group,  
8 an N-terminal alkyl-type protecting group or a di or  
9 poly-peptide containing amino acids of the D- or L-  
10 configuration;

11                   (b) A1 and A9 are either or both an aliphatic,  
12 cyclic or an aromatic amino acid residue of the D- or  
13 L-configuration;

14                   (c) B is an aliphatic, cyclic or aromatic amino  
15 acid residue of the D- or L- configuration;

16                   (d) C is an aliphatic, cyclic or aromatic amino  
17 acid residue of the D- or L- configuration;

18                   (e) D is an aliphatic, cyclic or aromatic amino  
19 acid residue of the D- or L-configuration;

20                   (f) W is an aliphatic, cyclic or aromatic amino  
21 acid residue of the D- or L- configuration;

22                   (g) X is an aliphatic, cyclic or aromatic amino  
23 acid residu of the D- or L- configuration;

24                   (h) Y is a D-aromatic amino acid residue or a  
25 substituted D-aromatic amino acid r sidue;

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26 (i) Z is an aliphatic, cyclic or aromatic amino  
27 acid residue of the D- or L- configuration; and

28 (j) Cn is a hydroxyl group or a C-terminal extension  
29 such as an amide, alkoxy group, an acidic, basic or neutral  
30 aliphatic, cyclic or aromatic amino acid residue of the D- or  
31 L-configuration or a peptide consisting of acidic, basic or  
32 neutral, aliphatic, cyclic or aromatic amino acid residues of  
33 the D- or L- configuration.

1 2. The modified bradykinin type peptide of claim 1  
2 wherein Y is a D-hydrophobic amino acid residue.

1 3. The modified bradykinin type peptide of claim 2  
2 wherein Al is a D-hydrophobic amino acid residue.

1 4. The modified bradykinin type peptide of claim 3  
2 wherein Y and Al are selected from the group consisting  
3 essentially of DPhe, DThi, DPAl, DNAl, CDF, DArg, DTry, FDF,  
4 DTrp, DVal and DIle.

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1           5. The modified bradykinin type peptide of claim 4  
2 wherein B, C and D are aliphatic, cyclic or aromatic amino  
3 acids of the D- or L- configuration and A9 is Arg and Cn is a  
4 hydroxyl group.

1           6. The modified bradykinin type peptide of claim 5  
2 wherein N is a hydrogen, Arg, Lys-Lys and Nal.

1           7. The modified bradykinin type peptide of claim 6  
2 having the formula  
3           DNal-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg.

1           8. The modified bradykinin type peptide of claim 6  
2 having the formula  
3           DArg-DNal-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg.

1           9. The modified bradykinin type peptide of claim 6  
2 having the formula  
3           Lys-Lys-DNal-Pro-Pro-Gly-Thi-S r-DPhe-Thi-Arg.

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1                   10. The modified bradykinin type peptide of claim 6  
2   having the formula

3                   DNal-DNal-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg.

1                   11. The modified bradykinin type peptide of claim 6  
2   having the formula

3                   DArg-DNal-Hyp-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg.

1                   12. The modified bradykinin type peptide of claim 6  
2   having the formula

3                   DArg-DNal-Pro-Hyp-Gly-Thi-Ser-DPhe-Phe-Arg.

1                   13. The modified bradykinin type peptide of claim 6  
2   having the formula

3                   DArg-DNal-Pro-Pro-Gly-Phe-Ser-DThi-Phe-Arg.

1                   14. The modified bradykinin type peptide of claim 6  
2   having the formula

3                   CDF-Pro-Pro-Gly-Phe-Ser-DPhe-Ph -Arg.

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1           15. The modified bradykinin type peptide of claim 6  
2   having the formula  
3           DArg-CDF-Pro-Hyp-Gly-Phe-Ser-DPhe-DPhe-Arg.

1           16. The modified bradykinin type peptide of claim 6  
2   having the formula  
3           DArg-Pro-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg.

1           17. The modified bradykinin type peptide of claim 6  
2   having the formula  
3           DArg-FDF-Pro-Hyp-Gly-Thi-Ser-DPhe-Phe-Arg.

1           18. The process of modifying the bradykinin type  
2   peptide of the formula of claim 1 or 5 wherein either or both  
3   B and C are changed to confer tissue selectivity.

1           19. The modified bradykinin type peptide of claim 1  
2   wherein Cn is selected from the group consisting essentially  
3   of NH<sub>2</sub> and Ile-Tyr.



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1           20. The modified bradykinin type peptide of claim  
2           19 wherein N is selected from the group consisting essentially  
3           of H, Ac, Lys-Lys and Arg.

1           21. The modified bradykinin type peptide of claim  
2           20 wherein Y is a D-hydrophobic amino acid residue.

1           22. The modified bradykinin type peptide of claim  
2           21 having the formula  
3           Ac-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg.

1           23. The modified bradykinin type peptide of claim  
2           21 having the formula  
3           Lys-Lys-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg.

1           24. The modified bradykinin type peptide of claim  
2           21 having the formula  
3           DArg-Arg-Pro-Pro-Gly-Phe-Ser-DPhe-Phe-Arg-Ile-Tyr.

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1                    25. The modified bradykinin type peptide of claim  
2       21 having the formula  
3                    DArg-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg-Ile-Tyr.

1                    26. The modified bradykinin type peptide of claim  
2                    21 having the formula  
3                    DArg-Arg-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg-Ile-Tyr.

1            27. The modified bradykinin type peptide of claim  
2            21 wherein A9 is a D aliphatic amino acid residue.

1                    28. The modified bradykinin type peptide of claim  
2                    27 having the formula  
3                    DArg-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg.

1                    29. The modified bradykinin type peptide of claim  
2                    27 having the formula  
3                    Ac-Arg-Pro-Pro-Gly-Thi-Ser-DPhe-Thi-Arg.

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1                   30. The modified bradykinin type peptide of claim  
2   21 wherein A1 is Nal.

1                   31. The modified bradykinin type peptide of claim  
2   30 having the formula  
3       DArg-DNal-Pro-Hyp-Gly-Thi-Ser-DPhe-Thi-Arg-DIle-Tyr.

1                   32. The modified bradykinin type peptide of claim  
2   21 wherein A1 and A9 are selected from the group consisting  
3   essentially of Arg, Nal, and Phe.

1                   33. The modified bradykinin type peptide of claim  
2   32 having the formula  
3       DNal-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Phe.

1                   34. The modified bradykinin type peptide of claim  
2   32 having the formula  
3       DArg-DArg-Pro-Hyp-Gly-Phe-Ser-DPhe-Phe-Phe.

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1           35. The pharmaceutical composition consisting  
2 essentially of the compounds as defined in claim 1 wherein  
3 enhanced resistance to enzymatic degradation is achieved by  
4 modifying the moiety A.

1           36. The pharmaceutical composition consisting  
2 essentially of the compounds as defined in claim 35 possessing  
3 enhanced resistance to enzymatic degradation wherein A is  
4 D-Arg, D-Lys, Lys-Lys, L-Thi, Met-Lys or Gly-Arg-Met-Lys and  
5 the pharmaceutically acceptable salts thereof in combination  
6 with a pharmaceutically acceptable carrier.

1           37. The pharmaceutical composition consisting  
2 essentially of the compounds as defined in claim 1 wherein  
3 enhanced tissue selectivity is achieved by modifying the  
4 moiety B alone or in combination with a modification of the  
5 moiety C.

1           38. The pharmaceutical composition consisting  
2 essentially of the compounds as defined in claim 37 possessing  
3 enhanced tissue selectivity wherein B alone, C alone or a  
4 combination thereof are selected from a group comprising  
5 L-Pro, L-hydroxyproline, APro, D-valine, L-valine,  
6 alpha-ammoisobutyric acid (Aib), L-Ala, D-Ala, Sar, D-Gly,  
7 L-Gly and the pharmaceutically acceptable salts thereof in  
8 combination with a pharmaceutically acceptable carrier.

1           39. The pharmaceutical composition consisting  
2 essentially of the compounds as defined in claim 1 wherein  
3 enhanced potency is achieved by modifying the moiety W, X or Z  
4 alone or in combination.

1           40. The pharmaceutical composition consisting  
2 essentially of the compound as defined in claim 39 possessing  
3 enhanced potency wherein W alone or Z alone or in combination  
4 are selected from a group comprising Phe, O-Methyl-Tyr(OMT),  
5 p-Chloro-Phe(CLF), p-Nitro-Phe(PNF), beta-2-naphthyl-Ala(Nal),  
6 Tyr, Thi, Pal, and the pharmaceutically acceptable salts  
7 ther of in combinati n with a pharmac utically acceptable  
8 carrier.

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1                   41. The pharmaceutical preparation for treating  
2 local pain and inflammation from burns, wounds, cuts, rashes  
3 and other such trauma and pathologic conditions caused by the  
4 production of bradykinin or related kinins by the animal which  
5 comprises an effective amount of the compound of claim 1  
6 sufficient to antagonize bradykinin and a suitable  
7 pharmaceutical carrier.

1                   42. A process for treating local pain and  
2 inflammation which comprises administering an effective  
3 amount of the pharmaceutical preparation of claim 41 to a  
4 host.

1                   43. The pharmaceutical preparation for treating  
2 pain, inflammation and swelling from bites, stings or other  
3 injection of bradykinin or related kinins which comprises an  
4 effective amount of the compound of claim 1 sufficient to  
5 antagonize bradykinin and a suitable pharmaceutical carrier.

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1           44. A process for treating local pain, inflammation  
2 and swelling which comprises administering an effective amount  
3 of the pharmaceutical preparation of claim 43 to a host.

1           45. The pharmaceutical preparation for treating  
2 rhinitis and other such trauma and pathologic conditions  
3 comprising an effective amount of the compound of claim 1 and  
4 a suitable pharmaceutical carrier.

1           46. A process for treating rhinitis which comprises  
2 administering an effective amount of the compound of claim 45  
3 to a host.

1           47. The pharmaceutical preparation for treating low  
2 blood pressure and other such trauma and pathologic conditions  
3 caused by the production of bradykinin or related kinins by  
4 the animal which comprises an effective amount of the compound  
5 of claim 1 and a suitable pharmaceutical carrier.

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II                    48. A process for treating low blood pressure which  
2 comprises administering an effective amount of the  
3 pharmaceutical preparation of claim 47 to a host.

I                    49. The pharmaceutical preparation for treating  
2 asthma and other such trauma and pathologic conditions caused  
3 by the production of bradykinin or related kinins by the  
4 animal which comprises an effective amount of the compound of  
5 claim 1 and a suitable pharmaceutical carrier.

I                    50. A process for treating asthma which comprises  
2 administering an effective amount of the pharmaceutical  
3 preparation of claim 49 to a host.

I                    51. The pharmaceutical preparation for treating  
2 arthritis and other such trauma and pathologic conditions  
3 caused by the production of bradykinin or related kinins by  
4 the animal which comprises an effective amount of the compound  
5 of claim 1 and a suitable pharmaceutical carrier.



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1           52. A process for treating arthritis which  
2 comprises administering an effective amount of the  
3 pharmaceutical preparation of claim 51 to a host.

1           53. The pharmaceutical preparation for treating  
2 diarrhea and other such trauma and pathologic conditions  
3 caused by the production of bradykinin or related kinins by  
4 the animal which comprises an effective amount of the compound  
5 of claim 1 and a suitable pharmaceutical carrier.

1           54. A process for treating diarrhea which comprises  
2 administering an effective amount of the pharmaceutical  
3 preparation of claim 53 to a host.

1           55. The pharmaceutical preparation for treating  
2 irritable bowel syndrome and inflammatory bowel disease and  
3 other such trauma and pathologic conditions caused by the  
4 production of bradykinin or related kinins by the animal which  
5 comprises an effective amount of the compound of claim 1 and a  
6 suitable pharmaceutical carrier.

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1           56. A process for treating irritable bowel syndrome  
2           and inflammatory bowel disease which comprises administering  
3           an effective amount of the pharmaceutical preparation of claim  
4           55 to a host.

1           57. The pharmaceutical preparation for treating  
2           carcinoid syndrome and other such trauma and pathologic  
3           conditions caused by the production of bradykinin or related  
4           kinins by the animal which comprises an effective amount of  
5           the compound of claim 1 and a suitable pharmaceutical carrier.

1           58. A process for treating carcinoid syndrome which  
2           comprises administering an effective amount of the  
3           pharmaceutical preparation of claim 57 to a host.

1           59. The pharmaceutical preparation for treating  
2           pain associated with angina and other such trauma and  
3           pathologic conditions caused by the production of bradykinin  
4           or related kinins by the animal which comprises an effective  
5           amount of the compound of claim 1 and a suitable  
6           pharmaceutical carrier.

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1           60. A process for treating pain associated with  
2   angina which comprises administering an effective amount of  
3   the pharmaceutical preparation of claim 59 to a host.

1           61. The pharmaceutical preparation for treating  
2   pain and inflammation caused by the production of bradykinin  
3   or related kinins by the animal which comprises an effective  
4   amount of the compound of claim 1 and a suitable  
5   pharmaceutical carrier.

1           62. A process for treating pain and inflammation  
2   which comprises administering an effective amount of the  
3   pharmaceutical preparation of claim 61 to a host.

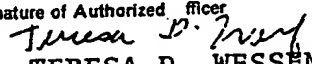
1           63. The pharmaceutical preparation for treating  
2   anaphylactic and septic shock and other such trauma and  
3   pathologic conditions caused by bradykinin or related kinins  
4   in the animal which comprises an effective amount of the  
5   compound of claim 1 with a suitable pharmaceutical carrier.

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- 1           64. A process for treating anaphylactic and septic
- 2           shock which comprises administering an effective amount of the
- 3           pharmaceutical preparation of claim 63 to a host.

# INTERNATIONAL SEARCH REPORT

International Application No. PCT/US88/02960

<b>I. CLASSIFICATION OF SUBJECT MATTER</b> (If several classification symbols apply, indicate all) <sup>6</sup>		
According to International Patent Classification (IPC) or to both National Classification and IPC <b>INT. CL(4): A61K 37/42; C07K 7/18</b> <b>U.S. CL.: 514/14, 15; 530/314, 327, 328</b>		
<b>II. FIELDS SEARCHED</b>		
Minimum Documentation Searched <sup>7</sup>		
Classification System	Classification Symbols	
U.S.	514/14, 15, 803; 530/314, 327, 328	
Documentation Searched other than Minimum Documentation to the Extent that such Documents are Included in the Fields Searched <sup>8</sup>		
<b>III. DOCUMENTS CONSIDERED TO BE RELEVANT <sup>9</sup></b>		
Category <sup>10</sup>	Citation of Document, <sup>11</sup> with indication, where appropriate, of the relevant passages <sup>12</sup>	Relevant to Claim No. <sup>13</sup>
X	US, A, 3,364,193 (HEMPEL) 16 January 1968, see the entire document.	1
A	US, A, 4,399,124 (FAUVE) 16 August 1983, see the entire document.	1-64
X	Chemical Abstracts, Vol. 88, no. 7, issued 1978 (Columbus, Ohio, USA), Moore, Stanley 'Synthesis of analogs of bradykinin with replacement of the arginine residues by 4-guanidino-phenyl-L-alanine, p. 579, col. 1, abst. no. 51156e, J. Chem. Soc., Perkin Trans 1 1977, (18) 2025-30 (Eng).	1
<div style="display: flex; justify-content: space-between;"> <div style="width: 45%;"> <p><sup>14</sup> Special categories of cited documents: <sup>15</sup></p> <p>"A" document defining the general state of the art which is not considered to be of particular relevance</p> <p>"E" earlier document but published on or after the international filing date</p> <p>"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)</p> <p>"O" document referring to an oral disclosure, use, exhibition or other means</p> <p>"P" document published prior to the international filing date but later than the priority date claimed</p> </div> <div style="width: 45%;"> <p>"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention</p> <p>"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step</p> <p>"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.</p> <p>"Δ" document member of the same patent family</p> </div> </div>		
<b>IV. CERTIFICATION</b>		
Date of the Actual Completion of the International Search		Date of Mailing of this International Search Report
15 November 1988		09 JAN 1989
International Searching Authority		Signature of Authorized Officer
ISA/US		 TERESA D. WESSENDORF

## III. D CUMENTS CONSIDERED TO BE RELEVANT (CONTINUED FROM THE SECOND SHEET)

Category *	Citation of Document, with indication, where appropriate, of the relevant passages	Relevant to Claim No
X	Chemical Abstracts, Vol. 91, no. 1, issued 1979 (Columbus, Ohio, USA), Skaric, V. ' $\beta$ -Alanine, $\delta$ -aminobutyric acid analog of bradykinin', page 512, col. 1, abs. no. 54495, Croat Chem. Acta. 1978 (Pub. 1979). 51(4), 347-51 (Eng).	1-3
A	Chemical Abstracts, Vol. 91, no. 21, issued 1979 (Columbus, Ohio, USA), St. Pierre, S. 'Synthesis of peptides by the solid-phase method. IV. Des-Arg <sup>9</sup> -bradykinin and analogs', page 95, col. 2, abs. no. 168871n, Can. J. Biochem. 1979, 57 (8), 1084-9 (Eng).	1-64
X	Chemical Abstracts, Vol. 92, no. 21, issued 1980 (Columbus, Ohio, USA), Filatova, M. P. 'Synthesis of new bradykinin analogs with a modified peptide chain', see page 701, col. 1, abs. no. 181624z, Bioorg. Khim., 1979, 5(10), 1445-54 (Russ).	1-3
A	Chemical Abstracts, Vol. 101, no. 11, issued 1983 (Columbus, Ohio, USA), Barabe, J. 'New agonist and antagonist analogs of bradykinin', see page 73 col. 1, abs. no. 84128n, Can. J. Physiol. Pharmacol. 1984, 62(6), 627-9 (Eng).	1-64
X	Chemical Abstracts, Vol. 103, no. 13, issued 1985 (Columbus, Ohio, USA), Vavrek, Raymond J. 'Competitive antagonist of bradykinin', see page 69, col. 2, abs. no. 98853y, Peptides (Fayetteville, N.Y.) 1985, 6(2), 161-4(Eng).	1-6,18, 21,35,36, 39,40,47, 48 and 60
X	Chemical Abstracts, Vol. 105, no. 21, issued 1986, (Columbus, Ohio, USA), Vavrek, Raymond J. 'Development and modification of competitive antagonists of bradykinin', see page 85, cols. 1 and 2, abs. no. 184127n, Pept. Struct. Funct., Proc. Am. Pept. Symp., 9th 1985, 655-8 (Eng)	1-6,18, 21,35,36, 39,40,47, 48 and 60